

Sequence Listing

SEQ ID NO.: 1: ESX cDNA ORF and deduced amino acid sequence
(See Figure 1) ??

SEQ ID NO.: 2: ESX cDNA sequence (5' untranslated + ORF + 3' untranslated) 1907 b.p.

cggccagataacctcagcgctacactggcggaactggatttctccgcctgcccgcctgcct
gccacagccggactccgccactccggtagcctcatggctgcaaccgtgagattagcaacat
tttagcaactacttcagtgcgtacagctcgaggactccacccctggcctctgttcccc
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gagggtacagagaaggccagctgggtggggAACAGCCAGTTCTGGTGAAGACGCAAGGT
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gtgtgggacccttaggatggggctccagctccctctgtgaatggaggcagagacc
tccaataaaagtgccttctggctttctaaaaaaaaaaaaaaaaaaaaaa

SEQ ID NO.: 3: Complete ESX deduced amino acid sequence (see Figure 1)

SEQ ID NO.: 4: First variable region (nucleotides 1-189 of Figure 1)

atggctgcaacctgtgagattagcaacattttagcaactacttcagtgcgtacagctc
ggaggactccaccctggcctctgttcccccctgctgccacccttggggccatgacttggta
ctgaccctgagcaaccccccagatgtcattggagggtacagagaaggccagctgggtggggaa
cag

SEQ ID NO.: 5: ~~Pointed~~ region (nucleotides 190-309 of Figure 1)

ccccagttctggtcgaagacgcagggtctggactggatcagctaccaagtggagaagaacaa
gtacgacgcagcgccattgacttctcacgtgtgacatggatggcgccaccctctgc

SEQ ID NO.: 6: Second variable region (nucleotides 210-561 of Figure 1)

aattgtgcccttggaggagctgcgtctggctttggcctctggggaccaactccatgccc
gctgcgagacccacttccagcttctgtatgagctcagttggatcattgagctgctggaga
aggatggcatggccttcaggaggccctagaccaggcccttgaccaggcagccccctt
gcccaggagctgctggacgacggtcagaagccagccctaccacccoggcagctgtggcgc
agga

SEQ ID NO.: 7: Deduced amino acid sequence for second variable region (amino acids 104-187 of Figure 1)

asn-cys-alanine . . . glycine-alanine-glycine

SEQ ID NO.: 8: Serine-rich region (nucleotides 562-714 of Figure 1)

gccccctccctggcagctctgacgtctccaccgcagggactggtgcttcggagctccca
ctcctcagactcccggtggaagtgacgtggacctggatcccactgtggcaagctctccccca
gcgatggtttcgtgactgcaagaagggg

SEQ ID NO.: 9: Third variable region (nucleotides 715-819 of Figure 1)

gatcccaagcacggaaagcgaaaacgaggccggcccgaaagctgagcaaagagtactggga
ctgtctcgaggcaagaagagcaagcacgcgcccagaggcacc

SEQ ID NO.: 10: Ets DNA Binding domain (nucleotides 820-1062 of Figure 1)

cacctgtggagttcatccggacatcctcatccacccggagctcaacgaggccctcatgaa
gtgggagaatcgcatgaaggcgcttcaagttcctgcgcctcgaggctgtggcccaactat
ggggccaaaagaaaaagaacagacaacatgac tac gaga agctgagccggccatgaggta
tactacaaacggagatcctggAACGGTGGATGGCCGGCAGTCGTCTACAAGTT

SEQ ID NO.: 11: Fourth variable region (nucleotides 1063-1113 of Figure 1)

ggcaaaaactcaagcggttggaggaggaaagagggttctccagagtcggAAC

SEQ ID NO.: 12: C-terminal 16 amino acids (amino acids 356-371 of Figure 1)

lys-asn-ser . . . ser-arg-asn

SEQ ID NO: 13 5'ESX-DBD

5'-CCGGGACATCCTCA TCCACCC-3'

SEQ ID NO: 14 3' ESX-DBD

5'-GTACCTCATGGCCCGGCTCAG-3' (SEQ ID NO. 14)).

SEQ ID NO: 15 Mouse ESX genomic sequence.

1 GGATCCTTCC AAGGCAGTGA CCTCACCCAA TTCTTTCTCA CTTTTCTCCT
51 CCATTAACT GTGGACGGAA TCAATACTCA GGGGGATGCG CTAGCTCTAA
101 GATTTCTGCA GCTTTGCCTC TCCTGAGCGG AAGCCCCGTG AAGGCAAGGG
151 AGCTAGCTGA TGGACTCTTT GTGGTCTTCT TCCTCTTGCG TCTGGAGACC
201 CAACCAGGTG TTCTTAGGGG AAGGAGCAG TGAGTAGCCA AGAGGCTAAA
251 AGCTGGTTCT CCCACATTCC AGGGTAAGTG ACTGGGTAGA GGGTGTGTCT
301 GCCTCAGGCT GCTTGGAGGA GGTCCCCTGA AGGGCCATGA GAAAATCCTA
351 CCCAGAGCCC TTGGTTTCC AGCAGCCCTC CACCTAGAGG AAAGGAGCCT
401 GTCGTTCTGA AGATGAAGAG TGGAGCCTAT GGGGGTGGGC AGATTGTGTC
451 CTGGGACAAT GGGGTACCTA GAAGAGAAAG GAATCTCCTT TCGTTTGAGG
501 TCTACCTGGG GGTCGTGTGT CTGTAAATGG GGTGGAGAGA GGAGAAGACA
551 CAGATCTTAT AACGTAGATG CAGGAAATGC TGACAGTTCA GTGTAGAGAA
601 CTTACTCAAT TCATATAGCC TCCAAAGCTA TCTCCTCAGG CAACGCAAAA
651 CAAACCAGTT GGAGCCGCAA GACATCTAAT GGCTTATCGA GTCCCACACC
701 CTCGATTCTT TGCTAATTTC ATGGTTTGC TTTTGAGACA ATCTACTGTA
751 GCCTAAGATA GCCCCAAACT CAAATGTAGC TGAGGGCTGAC TGACCCCTGAG
801 CTCTGGAATT CCAGACACAT GCATATCTT TGCTAGGCAA TAATCGCTCT
851 ACCAGCTGTA CTCCCACATT CCAGGGTAAG TGACTGGAAT TCTCACTTAC
901 TATATCCCTT TAAAAATTCC CTGAGTGGGA TGGTTGTAGC CAGAGGGAAA
951 AGGCACCAAC AACTGCTTGT CACTTTCCAA ATTTGGTAGC CTGAACAAAC
1001 CACTTATCAA GACAACAAC ATATATCATT TCTTTCTTC TCTCTCTCTC
1051 TCTCTCTCTC TCTCTCTCTC TCTCTCTCTC TCTCTCTCTC TCTCTCTTNN
1101 GAAAGAGTCT CACTACTATG TAGCCCTTGA TAACCTAGAA CTCACTATGT
1151 AGTCCAGGCT TGGCCTTCAG CTCGCAGAGG TCCACTTGCC TTGGGAGTTG
1201 AGAGATTAAA GGGATGCATC TCCACATGTG TCCAACAGTG CTTTTTAAAA

1251 ATATTTTAA AACCATGCTT ACAGCCAGGC ATAGTGGCG TGCCCTTAAT
1301 CCCAGTACTG GGGAGGCAGA GGTAGGTAGA GTTCTGAGTT GGAGGCTAGC
1351 CACATAGTAA GTCCCAGGAT AGCTAGAACT ATGTAAAGAC CATGTCTCAA
1401 AAAAGATGCA CACACACATA TACACACACA CGTTGTATG TGTTTGTAA
1451 GTGTGTATGT GTGTGTACAC TTGCACATAA AGGTCAGAGT ACCACATTAC
1501 AGGAGTCAGT TTTCTCCTTT TATCATGTAT GGATGGAACA CGGGTCCATC
1551 CATAGCATCC TTAGCAGCAG GTATCCTTAT CCACTGAGCT ATCTCAGCAG
1601 CCCCCACATTG CTTATTGGAT GTTTTGGAAT GAGGATAGTT ATATTAAAAAA
1651 GGTTTCTGGT GTTGGTCTGG GTAGTTACCC TTTAACCCAT CTCTAGAGCC
1701 TGTCTCTTGA GTTTGAGGCC AGCCTGGTAT ATGTAGCTAG ACAAAAGTTTC
1751 AAAATGAAC AGAACCTGG GACTAGAAC CATTGTAGA ATGCTTGCAT
1801 AAGAAGCTCT GGGTTCAACT TCCTGCATCT CCAGAGGGAT TTTGTTCTGT
1851 AGTTTAGTT TTTCAAGACA GAGTTCTCT GTGTAGCCCT GGCTGTCTG
1901 GAACTCACTC TGTAGACAAG GCTGGCCTCG AACTCAGAAA TCCTTCTACC
1951 TCTACTTCAG GAGTGCTGGG ATTAAAGATG TGCGCTGCC TCCTCCACCC
2001 CAATTGTTT TTGTTTTTA AGGGCCCCGG TAAACAGTAA ATTAACATGT
2051 GCATCCTGTT TGTCTTGTA ATGACTCAA TGTTGGCTT CTGACCACTA
2101 GAGGGCAGCA GGCAGATACT AATGGACTGG GCGGAGAGAA GGGTAATCAG
2151 GAGCAGACCA GACTCGCGGA TAAACCAAAC AGCACCGCCA GCCGACCCTA
2201 GGCAGGGAGA GCGCCACAGG CACCAAGGGGA AGACTTGAAG TAGTGTCTGA
2251 TCTCTACCGC TTCAGCAACC ATCGCGTTG GGTGGGCTCC AGACAGGCAA
2301 AGTGCAGCA AATGGTCCCT GTAGCTGACT AAACAGACTA TCAGACCCAA
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2401 GAATCCCAGC TTCTGGGTGT TGTGGAGGAA ACCCCTTAGC CTCGGTAACT

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2451 TTCACCAGGC CCTTCTTGTGTC TCTAGACATC TAGACAGTTG GAAGCATCAG
2501 TCTTGACCCA GCCACCGGTT CAGATTCTTT GCCTTGCTTT TTCTTCCCCA
2551 GTTCAGCCCT GGCCAGGCC CCAGGAAGAA TTTCCAGGGC CAGAGGGCAG
2601 CCTAAGGCAC AGATGCCAC CCCTGCAATG TTCCCGCCAC ATGCCAGTT
2651 CAGTACCCAG GGCCCAACCC CAGAGGGTGC GGAATGACAG ATTCTGACAA
2701 TCATTAAACC AGCCAGGCCT GATTTCCCAG CACCGCCCGT TAGGATATGG
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2851 CGGGGACGTA CGCCGAAGAC CTGGAGGGGA GGAGCTCCTG CTTTGCTCTA
2901 TTTAGAGCGG GTGGGGGCAG CGCCCTGGCC ACACTCATCA CTGCTACCTG
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3001 TGCTCACCCG CCTGCCACAC CGAACCTGA CACACCTCGG TACGGTCACA
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3251 AGAATTGTGTA TCATGCTCCC ACCCGCTTG AGATTATTT TATTTTCGG
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3351 GAGAGCATCC AACCTCAGCT TCCCCAAGTA GCTGGCTCTT GGTGGTGATG
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3701 CCTTTGGCAC TGAAGACTTG GTGTTGACCC TGAACAACCA ACAGATGACA
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3801 ATGATCTGAG AGGCTCTTAG ATGATAAAATG GACAGGGAGG AAGGGTATCC
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4001 GACAGATTAA GGGAGGATGA ACTTGAGAAC TAGCCATGTT GTGATTGTGG
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6901 AGAGAGGGCC AGCTCAGCAC ACTGGGGCTG GGAACCAATG CGAACCTCAG
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7101 GAGAGTCGGA ATTAAGGATC GGGGCTGGAC CCAGGACCTG ACTCAGGCAT
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7201 CCTTAACATG GATGTGTTCC CTGTGTTGCT GTAGAGAGGA AGAACCTGTT
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7301 GCCCTCTTGG AATTACAAGC CCCGGGTTTG AACCAAATTG TTGATAACT
7351 CTTCCAGCTG TGATTCCAGT TCCCTCCCGT CCCAACATGG ACTGCAAATG
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7451 GGCAGGAGAC TGCAGGGACG GAGGGGACAG GGTTGTGTCC TCGGTACTTC
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7551 GTGGATCAT CCCTAATTAA TGTGCTATAA ATATTCCAGG TGTATATAGA
7601 GAGCTATTTT TTCTAAAGCA TTTCCCTCC CTGCTCTTCT CCACTGAGTG
7651 CTGGTGGCCA GACTGATTTT TTTTTAGCC CCCCTAACTG GACCAGCGAG
7701 AAGTAGGGTG ATTCCAGGAC CCCCTCTTCC CCCAGAGGGG TCTCCTGGAT
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